### **Modify Query**

```
glutathione transferase (EC 2.5.1.18) 8, cytosolic - rat
XURT8C
       glutathione transferase (EC 2.5.1.18) class alpha chain Ya1 - rat
XURTG
       glutathione transferase (EC 2.5.1.18) pi - human
A37378
       glutathione transferase (EC 2.5.1.18) 7 - rat
XURTGP
       glutathione transferase (EC 2.5.1.18) 4 - rat
XURTG4
       glutathione transferase (EC 2.5.1.18) 1 - fruit fly (Drosophila
XUFF11
         melanogaster)
       glutathione transferase (EC 2.5.1.18) class I - house fly
A42045
       glutathione transferase (EC 2.5.1.18) - greenbottle fly (Lucilia
S43851
        glutathione transferase (EC 2.5.1.18) I - maize
XUZM1
        glutathione transferase (EC 2.5.1.18) III (version 1) - maize
XUZM31
        glutathione transferase (EC 2.5.1.18) III (version 2) - maize
XUZM32
       glutathione transferase (EC 2.5.1.18), fosfomycin-modifying -
A60635
          Escherichia coli plasmid pSU961 transposon Tn2921
       glutathione transferase (EC 2.5.1.18) yndN, fosfomycin-modifying -
E69890
          Bacillus subtilis
       glutathione transferase (EC 2.5.1.18), fosfomycin-modifying -
B48175
          Staphylococcus epidermidis
       glutathione transferase (EC 2.5.1.18) alpha (clone pGT41) - mouse
C28946
       glutathione transferase (EC 2.5.1.18) class alpha chain Ya1 - mouse
A27848
       glutathione transferase (EC 2.5.1.18) alpha-II, hepatic - rabbit
B41031
       glutathione transferase (EC 2.5.1.18) alpha-I, pulmonary - rabbit
A41031
       glutathione transferase (EC 2.5.1.18) Yc1 - rat
A26753
       glutathione transferase (EC 2.5.1.18) alpha-2 (clone GTH2) - human
S24330
       glutathione transferase (EC 2.5.1.18) alpha-2 (clone GTH2(+)) -
S77958
          human
        glutathione transferase (EC 2.5.1.18) 1 - rat (fragment)
A24735
       glutathione transferase (EC 2.5.1.18) 1-2 - rat (fragment)
B24735
        glutathione transferase (EC 2.5.1.18) a - guinea pig
JX0294
        glutathione transferase (EC 2.5.1.18) omega-1 chain - human
S29657
          (fragments)
       glutathione transferase (EC 2.5.1.18) omega-2 chain - human
S29658
          (fragments)
        glutathione transferase (EC 2.5.1.18) alpha-1 - chicken
S43431
        glutathione transferase (EC 2.5.1.18) alpha-2 - chicken (fragment)
S43432
        glutathione transferase (EC 2.5.1.18) isoenzyme 5.8 - human
S45687
          (fragments)
        glutathione transferase (EC 2.5.1.18) - rat (fragment)
S10555
        glutathione transferase (EC 2.5.1.18) alpha A5 - rat
A54858
        glutathione transferase (EC 2.5.1.18) 5.7 - mouse
S27234
        glutathione transferase (EC 2.5.1.18) - human
S20331
        glutathione transferase (EC 2.5.1.18) class alpha chain Ya3 - mouse
S24322
        glutathione transferase (EC 2.5.1.18) - chicken
S19734
        glutathione transferase (EC 2.5.1.18) alpha-Yx - rat (fragment)
S21278
        glutathione transferase (EC 2.5.1.18) GSTA2 - human
A56666
        glutathione transferase (EC 2.5.1.18), pancreatic - human
S45689
        glutathione transferase (EC 2.5.1.18), brain - human (fragments)
S45690
        glutathione transferase (EC 2.5.1.18) class alpha chain 10 - rat
S14355
          (fragments)
        glutathione transferase (EC 2.5.1.18), cardiac - human (fragments)
S45688
        glutathione transferase (EC 2.5.1.18) class alpha chain Ya2,
A26653
          hepatic - rat
```

\$29098 glutathione transferase (EC 2.5.1.18) class alpha A2 - golden hamster (fragments)

152381 glutathione transferase (EC 2.5.1.18) - human

152082 glutathione transferase (EC 2.5.1.18) A3 - human (fragment)

152097 glutathione S-transferase Ya subunit - rat (fragment)

165274 glutathione transferase (EC 2.5.1.18) class alpha A1 - golden hamster (fragments)

165274 glutathione S-transerase Ya subunit (put.) - rat (fragment)

165274 glutathione transferase (EC 2.5.1.18) Ya1, class alpha - mouse (fragments)

165274 glutathione transferase (EC 2.5.1.18) Ya1, class alpha - mouse (fragments)

165274 glutathione transferase (EC 2.5.1.18) Ya1, class alpha - mouse (fragments)

1752882 glutathione transferase (EC 2.5.1.18) / fatty-acyl-ethyl-ester synthase (EC 3.1.1.67) III, myocardial - human

isulice ⊬el.	(1) (1) (1)
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### **Modify Query**

### **Modify Query**

Field

#### **Query String**

1. Create the list of all entries containing:

1	All fields	dutathione transferace
	All licius	glutathione transferase

2. • refine, O add to, O remove from the list with entries containing:

	-
All fields	
All Holds	
	_

3. • refine, O add to, O remove from the list with entries containing:

All fields		
/ ut tioldo		

4. • refine, O add to, O remove from the list with entries containing:

All fields	

### **Modify Query**

```
KUHU
        ferroxidase (EC 1.16.3.1) precursor - human
        ferroxidase (EC 1.16.3.1) precursor - rat
A35210
        u-plasminogen activator (EC 3.4.21.73) precursor - human
UKHU
        u-plasminogen activator (EC 3.4.21.73) precursor - yellow baboon
UKBAY
UKPG
        u-plasminogen activator (EC 3.4.21.73) precursor - pig
JN0560
        u-plasminogen activator (EC 3.4.21.73) precursor - bovine
        u-plasminogen activator (EC 3.4.21.73) precursor - mouse
UKMS
S18932
        u-plasminogen activator (EC 3.4.21.73) precursor - rat
        u-plasminogen activator (EC 3.4.21.73) precursor - chicken
A35005
        t-plasminogen activator (EC 3.4.21.68) precursor - human
UKHUT
       t-plasminogen activator (EC 3.4.21.68) precursor - rat
A35029
A29941
        t-plasminogen activator (EC 3.4.21.68) precursor - mouse
        t-plasminogen activator (EC 3.4.21.68) precursor - false vampire
A34369
          bat (Megaderma lyra)
PLHU
        plasmin (EC 3.4.21.7) precursor - human
        plasmin (EC 3.4.21.7) precursor - pig (fragment)
PLPG
PLMS
        plasmin (EC 3.4.21.7) precursor - mouse
        plasmin (EC 3.4.21.7) precursor - bovine
PLBO
        basic proteinase inhibitor precursor - bovine
TIBO
        uterine plasmin/trypsin inhibitor - pig
<u> A55115</u>
       proteinase inhibitor (BPI-type) - horseshoe crab (Tachypleus
TIHCBP
          tridentatus)
        proteinase inhibitor (PSTI type), submandibular - dog
TIDGS
       proteinase inhibitor (PSTI type), submandibular - red fox
S03371
XSSMA
        subtilisin inhibitor precursor (PSTI type) - Streptomyces
          albogriseolus
XFSMF
        plasminostreptin (PSTI-type proteinase inhibitor) - Streptomyces
          "antifibrinolyticus"
        plasminogen activator inhibitor 1 precursor - human
ITHUP1
        plasminogen activator inhibitor 1 precursor - rat
A35032
806745 plasminogen activator inhibitor-1 precursor - bovine
       plasminogen activator inhibitor 1 homolog mrl - mouse
A34761
ITHUA2
       alpha-2-antiplasmin precursor - human
       urokinase-type plasminogen activator receptor precursor - bovine
JN0561
S06763
       calreticulin precursor - mouse
        fibrinogen alpha chain precursor, short splice form - human
FGHUA
FGBOB
        fibrinogen beta chain - bovine
TTHUN
        tetranectin precursor - human
        plasminogen-related protein precursor - human
JC1195
SGHU1V
        vitronectin precursor - human
        staphylokinase - phage S phi-C
PRSAK
        glyceraldehyde-3-phosphate dehydrogenase (EC 1.2.1.12) -
A42963
          Streptococcus sp.
        venom plasminogen activator (EC 3.4.21.-) precursor - Chinese green
A57290
          tree viper
JS0597
        t-plasminogen activator (EC 3.4.21.68) alpha-1 precursor - common
          vampire bat
        t-plasminogen activator (EC 3.4.21.68) alpha-2 precursor - common
JS0598
          vampire bat
       t-plasminogen activator (EC 3.4.21.68) gamma precursor - common
JS0600
          vampire bat
        t-plasminogen activator (EC 3.4.21.68) beta precursor - common
JS0599
          vampire bat
138098 t-plasminogen activator precursor, inactive endothelial splice form
```

- huma	
plasmin	precursor - lamprey (fragments)
plasmin	(EC 3.4.21.7) precursor - chicken (fragment)
plasmin	(EC 3.4.21.7) precursor - rat (fragment)
plasmin	(EC 3.4.21.7) precursor - goat (fragments)
plasmin	(EC 3.4.21.7) precursor - rabbit (fragment)
plasmin	(EC 3.4.21.7) precursor - dog (fragments)
Est	9× 3
Query	
	Modify Query
	• -
Field	Query String
	Query String fall entries containing:
te the list of elds efine, O a	f all entries containing:
te the list o	f all entries containing:  plasmin
te the list of lelds efine, O a lelds efine, O a	f all entries containing:  plasmin
te the list of lelds efine, O a lelds	f all entries containing:  plasmin  dd to, O remove from the list with entries containing:
te the list of leids  efine, O a leids  efine, O a leids	f all entries containing:  plasmin  dd to, O remove from the list with entries containing:
	plasmin plasmin plasmin plasmin

### **Modify Query**

4 entries	were found	
335711	tissue kal macaque	likrein (EC 3.4.21.35) precursor, prostatic - rhesus
541212	prostate-s	pecific serine proteinase (EC 3.4.21) - human
A32297	(fragmen	t) pecific serine proteinase (EC 3.4.21) precursor - human
A56881	prostate-s	pecific membrane antigen - human
		·
Modify	Query	
		Modify Query
	Field	Query String
1. Creat	e the list of all	l entries containing:
All fie	elds	prostate specific antigen
k		
2. • re	fine, O add 1	to, O remove from the list with entries containing:
Allfie	elds <b>S</b>	
3. • re	fine, O add t	to, O remove from the list with entries containing:
All fie	elds <b>a</b>	
	- 0	
4. • re	fine, O add i	to, O remove from the list with entries containing:
All fi	elds	
-		2016/201
	1	

### **Modify Query**

```
cerevisin (EC 3.4.21.48) precursor - yeast (Saccharomyces
A29358
          cerevisiae)
        subtilisin (EC 3.4.21.62) Carlsberg precursor - Bacillus
SUBSCL
          licheniformis
        subtilisin (EC 3.4.21.62) DY - Bacillus subtilis (strain DY)
SUBSD
        subtilisin (EC 3.4.21.62) BPN' precursor - Bacillus
SUBSN
          amyloliquefaciens
        subtilisin (EC 3.4.21.62) amylosacchariticus precursor - Bacillus
SUBSS
          subtilis
        subtilisin (EC 3.4.21.62) E precursor - Bacillus subtilis
SUBSI
       thermitase (EC 3.4.21.66) - Thermoactinomyces vulgaris
SUMYTV
       subtilisin (EC 3.4.21.62) 1 precursor - Bacillus sp. (strain TA39)
S23407
       subtilisin (EC 3.4.21.62) precursor - Bacillus sp. (strain TA41)
S25835
       endopeptidase K (EC 3.4.21.64) precursor - imperfect fungus
SUTIKA
          (Tritirachium album)
       cuticle-degrading proteinase (EC 3.4.21.-) precursor - imperfect
S22387
          fungus (Metarhizium anisopliae)
      alkaline proteinase (EC 3.4.21.-) precursor - fungus (Fusarium sp.)
JC2142
          (strain S-19-5)
       serine proteinase (EC 3.4.21.-) isp6 - fission yeast
S45493
          (Schizosaccharomyces pombe)
       alkaline proteinase (EC 3.4.21.-) precursor - fungus (Acremonium
JU0332
          chrysogenum)
        serine proteinase (EC 3.4.21.-) prbl precursor - fungus
S32905
          (Trichoderma harzianum)
        oryzin (EC 3.4.21.63) precursor - Aspergillus oryzae
SUASO
        serine proteinase alp (EC 3.4.21.-) - Aspergillus fumigatus
S22184
        alkaline serine proteinase (EC 3.4.21.-) precursor - yeast
A26955
          (Yarrowia lipolytica)
        subtilisin-like proteinase (EC 3.4.21.-) YSP3 precursor - yeast
S61986
          (Saccharomyces cerevisiae)
        hypothetical protein YCR045c - yeast (Saccharomyces cerevisiae)
S19458
        serine proteinase (EC 3.4.21.-) precursor - Aspergillus niger
JU0146
        alkaline proteinase (EC 3.4.21.-) A precursor - Vibrio
JS0173
          alginolyticus
        aqualysin (EC 3.4.21.-) I precursor - Thermus aquaticus
A35742
        subtilisin-like proteinase ag12 (EC 3.4.21.-) - Alnus glutinosa
S52769
        cucumisin (EC 3.4.21.25) precursor - muskmelon
A55800
C41335
        microbial serine proteinase (EC 3.4.21.-), intracellular - Bacillus
          polymyxa
        alkaline proteinase (EC 3.4.21.-), intracellular - Bacillus sp.
S27501
          (strain 221)
        serine proteinase (EC 3.4.21.-) epr precursor - Bacillus subtilis
SUBSMP
        serine proteinase (EC 3.4.21.-) precursor - Serratia marcescens
SUSEMM
          (strain RH1)
        serine proteinase (EC 3.4.21.-) precursor, cell-envelope-associated
B44858
          - Lactobacillus paracasei (strain NCDO 151)
        serotype-specific antigen (EC 3.4.21.-) precursor - Pasteurella
SWQPA1
          haemolytica (strain A1)
        streptococcal C5a peptidase (EC 3.4.21.-) precursor - Streptococcus
A35066
          pyogenes
        serine proteinase, calcium-dependent (EC 3.4.21.-) - Anabaena
PRAISB
          variabilis
        56K serine proteinase (EC 3.4.21.-) - ictalurid herpesvirus 1
PRBEI1
```

(strain aub furin (EC 3.4.21.75) precursor - human KXHUF furin (EC 3.4.21.75) precursor - mouse KXMSF furin (EC 3.4.21.75) precursor - rat KXRTF prohormone-processing proteinase (EC 3.4.21.-) PC2 precursor -KXHUC2 prohormone-processing proteinase (EC 3.4.21.-) PC2 precursor -KXMSC2 mouse prohormone-processing proteinase (EC 3.4.21.-) PC2 precursor - rat KXRTC2 prohormone-processing proteinase (EC 3.4.21.-) PC1/PC3 precursor -KXHUC1 human prohormone-processing proteinase (EC 3.4.21.-) PC1 precursor - rat KXRTC1 prohormone-processing proteinase (EC 3.4.21.-) PC1 precursor -KXMSC1 kexin (EC 3.4.21.61) 2 precursor - yeast (Saccharomyces cerevisiae) KXBY kexin (EC 3.4.21.61) 1 precursor - yeast (Kluyveromyces marxianus S01013 var. lactis) serine proteinase (EC 3.4.21.-) PACE4 precursor - human A39490 subtilisin-like proprotein convertase (EC 3.4.21.-) PC5 precursor -A48225 subtilisin-like proprotein convertase (EC 3.4.21.-) PACE4D - human JC2192 subtilisin-like proprotein convertase (EC 3.4.21.-) PACE4C - human JC2191 proteinase inhibitor (PSTI type), submandibular - dog TIDGS

Prince Lieu

### **Modify Query**

#### **Modify Query**

Field

#### **Query String**

1. Create the list of all entries containing:

All fields subtilisin

2. • refine. O add to, O remove from the list with entries containing:

All fields

3. • refine, O add to, O remove from the list with entries containing:

All fields

4. • refine, O add to, O remove from the list with entries containing:

All fields

2 0 0 1111

### **Modify Query**

```
6-phospho-beta-galactosidase (EC 3.2.1.85) - Lactococcus lactis
GLSOPL
       beta-galactosidase (EC 3.2.1.23) - Staphylococcus aureus
A27233
        6-phospho-beta-galactosidase (EC 3.2.1.85) - Lactobacillus casei
A29897
       beta-galactosidase (EC 3.2.1.23) lacZ - Escherichia coli
GBEC
       beta-galactosidase (EC 3.2.1.23) alpha chain - Escherichia coli
GBECE
JC1266 beta-galactosidase (EC 3.2.1.23) - yeast (Kluyveromyces marxianus
          var. lactis)
B69649 beta-galactosidase lacA - Bacillus subtilis
A29836 beta-galactosidase (EC 3.2.1.23) I - Bacillus stearothermophilus
A69798 beta-galactosidase homolog yesZ - Bacillus subtilis
JE0395 phospho-beta-galactosidase I - Lactobacillus gasseri
       phospho-beta-galactosidase II - Lactobacillus gasseri
JE0396
       beta-glucosidase (EC 3.2.1.21) - Bifidobacterium breve
JC5137
       beta-galactosidase (EC 3.2.1.23) - Klebsiella pneumoniae
A24925
       beta-galactosidase (EC 3.2.1.23) - Thermoanaerobacterium
JU0275
          thermosulfurigenes
       beta-galactosidase - Thermotoga maritima (strain MSB8)
F72283
       beta-lactamase (EC 3.5.2.6) - Streptococcus thermophilus (strain
A49750
          "A054)
       beta-galactosidase (EC 3.2.1.23) - Lactobacillus delbrueckii subsp.
A30093
         bulgaricus
       beta-galactosidase (EC 3.2.1.23) - Arthrobacter sp.
I39697
       beta-D-galactosidase - Thermotoga maritima (strain MSB8)
E72391
       beta-galactosidase - Thermotoga maritima (strain MSB8)
H72283
       transcription repressor of beta-galactosidase gene lack - Bacillus
C69649
          subtilis
A32611
       beta-galactosidase (EC 3.2.1.23) precursor - human
       beta-galactosidase (EC 3.2.1.23) precursor - mouse
A37086
       beta-galactosidase (EC 3.2.1.23) precursor - Xanthomonas manihotis
A57249
       beta-galactosidase (EC 3.2.1.23) - Bacillus circulans
JC5618
S34735 beta-galactosidase (EC 3.2.1.23) - Agrobacterium radiobacter
       beta-galactosidase (EC 3.2.1.23) - Escherichia coli (fragment)
S23694
       beta-galactosidase (EC 3.2.1.23) beta chain - Escherichia coli
B65096
       beta-galactosidase alpha chain - Escherichia coli (fragment)
$59625
       beta-galactosidase - Bacillus megaterium
T30551
       beta-galactosidase - Bacillus megaterium
T30574
       beta-galactosidase (EC 3.2.1.23) - Clostridium acetobutylicum
A39405
       beta-galactosidase (EC 3.2.1.23) isozyme 12 - Arthrobacter sp.
A56263
          (strain B7)
       beta-galactosidase (EC 3.2.1.23) - Streptomyces coelicolor
T29434
       probable beta-galactosidase - Streptomyces coelicolor
T35944
        beta-galactosidase (EC 3.2.1.23) - Sulfolobus solfataricus
S06762
       beta-galactosidase (EC 3.2.1.23) - Sulfolobus solfataricus (strain
JQ0767
          MT-\tilde{4})
       beta-galactosidase homolog F20M13.150 - Arabidopsis thaliana
T05687
       beta-galactosidase homolog M4E13.70 - Arabidopsis thaliana
T05771
        probable beta-galactosidase (EC 3.2.1.23) - Arabidopsis thaliana
T04269
       probable beta-galactosidase (EC 3.2.1.23) F23E13.200 - Arabidopsis
T04600
          thaliana
       probable beta-galactosidase (EC 3.2.1.23) F24L7.5 - Arabidopsis
T00787
          thaliana
       beta-galactosidase (EC 3.2.1.23) - wild cabbage
S52393
       probable beta-galactosidase (EC 3.2.1.23) precursor - apple tree
T17002
```

T04340 T06590 S41889 S51494 B32688 JS0610	probable beta-galac arabinogal beta-galac	ctos ase (EC 3.2.1.23) II precursor tomato beta-galactosidase (EC 3.2.1.23) - tomato ctosidase (EC 3.2.1.23) - garden asparagus lactan endo-1,4-beta-galactosidase - Aspergillus aculeatus ctosidase-related protein - human ctosidase (EC 3.2.1.23) / glycosylceramidase (EC 3.2.1.62) or - rat	254.00
Modify			=->
		Modify Query	
	Field	Query String	
1. Creat	e the list of a	Il entries containing:	
All fi	elds	beta-galactosidase	
		to, O remove from the list with entries containing:	
All fi	elds		
3. • re	efine, O add	to, O remove from the list with entries containing:	
All fi	elds		
4. • re	efine, O add	to, O remove from the list with entries containing:	
All fi	elds		

### **Modify Query**

```
LZHU
        lysozyme (EC 3.2.1.17) c precursor - human
        lysozyme (EC 3.2.1.17) c - baboon
LZBA
        lysozyme (EC 3.2.1.17) c 2 precursor - bovine
LZBO
        lysozyme (EC 3.2.1.17) 1 precursor - rat
LZRT
LZCH
        lysozyme (EC 3.2.1.17) c precursor - chicken
        lysozyme (EC 3.2.1.17) c - California quail
LZQJEC
       lysozyme (EC 3.2.1.17) c - common bobwhite
LZQJEB
        lysozyme (EC 3.2.1.17) c precursor - Japanese quail
LZQJE
        lysozyme (EC 3.2.1.17) c precursor - ring-necked pheasant
LZFER
        lysozyme (EC 3.2.1.17) c precursor - turkey
LZTK
        lysozyme (EC 3.2.1.17) c - helmeted guineafowl
LZUH
        lysozyme (EC 3.2.1.17) c precursor - duck
LZDK
        lysozyme (EC 3.2.1.17) c III - duck
LZDK3
        lysozyme (EC 3.2.1.17) c - plain chachalaca
LZOVE
        lysozyme (EC 3.2.1.17) c - pigeon
LZPY
        lysozyme (EC 3.2.1.17) c precursor - cecropia moth (fragment)
LZWK
       lysozyme (EC 3.2.1.17) - tobacco hornworm
A40796
        lysozyme (EC 3.2.1.17) precursor - silkworm
JC4233
        alpha-lactalbumin precursor - human
LAHU
        alpha-lactalbumin - Arabian camel
LACM
        alpha-lactalbumin - horse
LAHO
LAPG
        alpha-lactalbumin precursor - pig
        alpha-lactalbumin precursor - bovine
LABO
        alpha-lactalbumin - bovine (tentative sequence)
LABOZ
LAGT
        alpha-lactalbumin precursor - goat
        alpha-lactalbumin precursor - guinea pig
LAGP
        alpha-lactalbumin precursor - rat
LART
        alpha-lactalbumin - rabbit
LARB
        alpha-lactalbumin - red-necked wallaby
LAKGAW
        lysozyme (EC 3.2.1.17) g - black swan
LZWSG
        lysozyme (EC 3.2.1.17) g - goose
LZGSG
        lysozyme (EC 3.2.1.17) g - ostrich
LZOSG
       lysozyme (EC 3.2.1.17), tail-associated - phage T4
G5BPT4
       lysozyme (EC 3.2.1.17) - phage T4
LZBPT4
       lysozyme (EC 3.2.1.17) - phage phi-29
WMBPP9
        lysozyme (EC 3.2.1.17) - phage PZA
WMBP15
        lysozyme (EC 3.2.1.17) - phage P22
LZBP22
        lysozyme (EC 3.2.1.17) - phage PA2
WMBPP2
        lysozyme (EC 3.2.1.17) - phage 21
lysozyme (EC 3.2.1.17) - fungus (Chalara sp.)
LZBP21
MUKAD
       lysozyme (EC 3.2.1.17) M1 precursor - Streptomyces globisporus
MUSMM1
       lysozyme (EC 3.2.1.17) - Lactobacillus delbrueckii subsp.
MUBPM1
          bulgaricus phage mv1
        N-acetylmuramoyl-L-alanine amidase (EC 3.5.1.28) - phage T7
MUBPA7
        N-acetylmuramoyl-L-alanine amidase (EC 3.5.1.28) - phage CP-1
MUBPCP
        N-acetylmuramoyl-L-alanine amidase (EC 3.5.1.28) - phage CP-9
MUBPC9
        N-acetylmuramoyl-L-alanine amidase (EC 3.5.1.28) - phage CP-7
MUBPC7
        chitinase (EC 3.2.1.14) / lysozyme (EC 3.2.1.17) PZ precursor,
S51591
          pathogenesis-related - common tobacco
        lysozyme (EC 3.2.1.17) - bare-faced crassow
JE0185
        lysozyme (EC 3.2.1.17) c - dog
S48642
        lysozyme (EC 3.2.1.17) I - Australian echidna
A45660
```

Modify Query		
Middly Quely	Modify Query	
Field	Query String	
1. Create the list of a	all entries containing:	
All fields	lysozyme	
2. • refine, O add	d to, O remove from the list with entries containing:	
All fields		
3. • refine, O add	d to, O remove from the list with entries containing:	
All fields		
4. • refine, O add	d to, O remove from the list with entries containing:	
All fields		
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No entries were found	
Modify Query	
	Modify Query
Field	Query String
1. Create the list of al	ll entries containing:
All fields	pyruvate kinase
2. • refine, O add	to, O remove from the list with entries containing:
All fields	
3. • refine, O add	to, O remove from the list with entries containing:
All fields	
4. • refine, o add	to, O remove from the list with entries containing:
All fields	
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TT 1 7/ ·	zetown.edu (17-Jan-2000)

# **Error Report**

Internal Error.



4 entries were found
A39019 glucose dehydrogenase (acceptor) (EC 1.1.99.10) - fruit fly (Drosophila melanogaster)
B39019 glucose dehydrogenase (acceptor) (EC 1.1.99.10) - fruit fly
(Drosophila pseudoobscura)  A35459 glucose oxidase (EC 1.1.3.4) precursor - Aspergillus niger
S14129 glucose oxidase (EC 1.1.3.4) - Aspergillus niger (fragment)
Modify Query
Modify Query
Field Query String
1. Create the list of all entries containing:
All fields glucose oxidase
2. • refine, O add to, O remove from the list with entries containing:
All fields
3. • refine, O add to, O remove from the list with entries containing:
All fields
4. • refine, O add to, O remove from the list with entries containing:
All fields
Subscribe )

# **Error Report**

Internal Error.

## **Modify Query**

DEHUG6	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - human
A56686	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - mouse
S01233	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) precursor - rat
DEFFG6	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - fruit fly
	(Drosophila melanogaster)
A56841	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - Japanese
	pufferfish
DEYCG6	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - Synechococcus
	sp. (strain PCC 7942)
S13744	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - yeast
	(Saccharomyces cerevisiae)
S31337	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - yeast
	(Kluyveromyces marxianus var. lactis)
S57785	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - alfalfa
S60287	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - potato
B37855	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - Zymomonas
20:000	mobilis
D64947	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - Escherichia
<u> </u>	coli
A39864	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - Leuconostoc
1155001	mesenteroides
A47740	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - fruit fly
A47740	(Drosophila melanogaster) (strain OK93)
S77348	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - Synechocystis
377340	sp. (strain PCC 6803)
S57485	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - Aspergillus
337403	niger
S54720	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - Aspergillus
554720	niger
S29381	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - yeast (Pichia
52 7501	jadinii)
S71245	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) (clone E5) -
011215	Arabidopsis thaliana (fragment)
s37053	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - Erwinia
557055	chrysanthemi
E64077	probable glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) -
E04077	Haemophilus influenzae (strain Rd KW20)
B69964	probable glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) yqjJ -
003304	Bacillus subtilis
C70179	probable glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - Lyme
C/01/3	disease spirochete
m10667	hypothetical protein B0035.5 - Caenorhabditis elegans
T18657	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) 1, cytosolic -
T14894	
m1 400C	parsley glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) 2, cytosolic -
T14896	
m1 4000	parsley glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49), chloroplast -
T14890	
D21210	parsley probable glucose-6-phosphate 1-dehydrogenase (zwf) - syphilis
D71319	•
m07275	spirochete
T07375	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49), chloroplast -
m02044	potato
T03244	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - common tobacco
T03740	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) TPG18 - common

	tobacco (PC 1 1 1 40) F3T6 22
T00659	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) F3I6.22 - Arabidopsis thaliana
T09090	<pre>glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) (clone O28FA38) spinach (fragment)</pre>
T09089	<pre>glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) (clone O30A5) - spinach plasmid pZL1 (fragment)</pre>
T09088	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - spinach
T07655	glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - soybean (fragment)
G72289	<pre>glucose-6-phosphate 1-dehydrogenase - Thermotoga maritima (strain MSB8)</pre>
C72103	glucose-6-p dehyrogenase - Chlamydia pneumoniae (strain CWL029)
B70538	probable glucose-6-phosphate 1-dehydrogenase - Mycobacterium tuberculosis (strain H37RV)
B70917	probable zwf2 protein - Mycobacterium tuberculosis (strain H37RV)
F71859	glucose-6-phosphate 1-dehydrogenase - Helicobacter pylori (strain J99)
E64657	probable glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - Helicobacter pylori (strain 26695)
B71546	<pre>probable glucose-6-phosphate dehyrogenase - Chlamydia trachomatis   (serotype D, strain UW3/Cx)</pre>
<u>H70916</u>	<pre>probable glucose-6-phosphate 1-dehydrogenase zwf2 - Mycobacterium tuberculosis (strain H37RV)</pre>
s7 <u>2775</u>	<pre>probable glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) isozyme gpdB - Mycobacterium leprae</pre>
F70 <u>127</u>	probable glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - Lyme disease spirochete
F64657	probable glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - Helicobacter pylori (strain 26695)
S18396	<pre>probable glucose-6-phosphate 1-dehydrogenase (EC 1.1.1.49) - Acetobacter hansenii (fragment)</pre>
B75377	<pre>glucose-6-phosphate 1-dehydrogenase - Deinococcus radiodurans   (strain R1)</pre>
<u>C75377</u>	probable glucose 6-phosphate dehydrogenase assembly protein OpcA -

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### **Modify Query**

Field

**Query String** 

1. Create the list of all entries containing:

	All fields glucose 6-phosphate dehydrogenase
2.	ego refine, O add to, O remove from the list with entries containing
	All fields
3.	ego refine, O add to, O remove from the list with entries containing
	All fields

All fields		
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### **Modify Query**

### 37 entries were found

Photinus-luciferin 4-monooxygenase (ATF-hydrolyzing) (EC 1.13.12.7)			
Photinus-luciferin 4-monooxygenase (ATP-hydrolyzing) (EC 1.13.12.7) Photinus-luciferin 4-monooxygenase (EC 1.14.14.3) alpha chain -	<u>\$2343</u> 7		
A26772 Photinus-luciferin 4-monoxygenase (ATP-hydrolyzing) (EC 1.13.12.7) - common eastern firefly 300574 alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) alpha chain - Vibrio fischeri 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) alpha chain - Photobacterium phosphoreum 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) alpha chain - Vibrio harveyi 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) alpha chain - Vibrio harveyi 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) alpha chain - Xenorhabdus luminescens 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) alpha chain - Xenorhabdus luminescens 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) alpha chain - Xenorhabdus luminescens 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) alpha chain - Xenorhabdus luminescens 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) alpha chain - Hashlight fish symbiont bacterium 31canal monooxygenase (FMN-linked) (EC 1.14.14.3) alpha chain - Photobacterium leiognathi 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) alpha chain - Vibrio cholerae (strain 14547) (fragment) 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) alpha chain - Vibrio cholerae (strain 1402) (fragment) 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) alpha chain - Vibrio cholerae (strain 1402) (fragment) 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) alpha chain - Vibrio cholerae (strain 1402) (fragment) 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain - Vibrio fischeri 31r954 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain - Photobacterium leiognathi 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain - Photobacterium leiognathi 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain - Photobacterium leiognathi 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain - Photobacterium leiognathi 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain - Photobacterium leiognathi 31kanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain - Xenorhabdus luminescens 31kanal monooxygenase (FMN-linke	<u>JS0181</u>		
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C35411 alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain - Xenorhabdus luminescens  C37898 alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain - Xenorhabdus luminescens  D38448 alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain - Xenorhabdus luminescens  A23866 alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain -	S18547	alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain -	
C37898 alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain - Xenorhabdus luminescens  D38448 alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain - Xenorhabdus luminescens  A23866 alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain -	C35411	alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain -	
D38448 alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain - Xenorhabdus luminescens A23866 alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain -	<u>C37898</u>	alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain -	
A23866 alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain -	D38448	alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain -	
	A23866	alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) beta chain -	

S33788	luciferase - Southern Russian firefly
557417	
S29355	luciferase - luminescent click beetle (Pyrophorus plagiophthalmus)
S29352	luciferase GR - luminescent click beetle (Pyrophorus
	plagiophthalmus)
S29354	luciferase YE - luminescent click beetle (Pyrophorus
	plagiophthalmus)
<u>\$293</u> 53	luciferase YG - luminescent click beetle (Pyrophorus
	plagiophthalmus)
S50144	luciferase - dinoflagellate (Crypthecodinium cohnii)
S53695	luciferase - dinoflagellate (Crypthecodinium cohnii)
A39374	Renilla-luciferin 2-monooxygenase (EC 1.13.12.5) - sea pansy
	(Renilla reniformis) (fragment)
G72605	probable luciferase APE1310 - Aeropyrum pernix (strain K1)

### **Modify Query**

Field	Query String
1. Create the list of all er	ntries containing:
All fields luci	ferase
2. • refine, O add to,	O remove from the list with entries containing:
All fields	
3. • refine, o add to,	O remove from the list with entries containing:
All fields	
4. • refine, O add to,	O remove from the list with entries containing:
All fields	